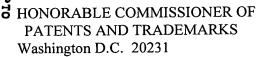








UTILITY PATENT APPLICATION UNDER 37 CFR 1.53(b)



Case Docket No. 114205.1200



Transmitted herewith for filing is the patent application of:

INVENTOR: Denise R. COOPER and Niketa A. PATEL

FOR: GLUCOSE REGULATED mRNA INSTABILITY ELEMENT

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[X]	124 pages of specification, claims, abstract
	Declaration & Power of Attorney
<u>[]</u>	Priority Claimed
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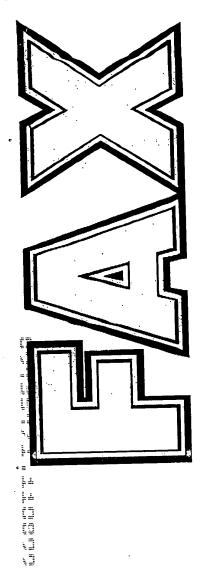
Respectfully submitted,

PEPPER, HAMILTON LLP

Gilberto M. Villacorta, Ph.D. Registration No. 34,038

by Corinne Pouliquen Registration No. 35,753

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APPENDIX 11

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INVENTION DISCLOSURE FORM DIVISION OF PATENTS AND LICENSING UNIVERSITY OF SOUTH FLORIDA **FAO 126**

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	INVENT		DISCLOSURE: Denise R. Cooper and Niketa A. Lato	
		TITLE: AS	sociate Professor and Research Post-doc	
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•	. 19	A.	What is the problem this invention addresses?	
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Title: Glucose-regulated mRNA instability element

2A. What is the problem this invention addresses?

Regulatable vectors for gene expression are often not reliable since they can be "leaky". They usually consist of systems where the gene of interest is cloned downstream of a minimal viral promoter fused to copies of the tetracycline operator. The promoters used with these repressor sequences are, however, not totally repressed in many cells. By engineering a construct that is also regulated at the posttranscriptional level, full repression of such a promoter could be achieved in a wider variety of cells. We propose that by inserting instability sequences down-stream of an inserted cDNA, genes can be further regulated using high extracellular concentrations of glucose or non-metabolized analogues. Both Tet and Retroviral Tet Systems (Clontech) show promise for regulating transcription of genes in cells. The ecdysone-inducible expression system derived from Drosophila is also commercially available (Invitrogen) and may have even lower basal activity in mammalian cells. Basal activity of the tetO.HCMC IE promoter is highly variable when tested in several cell lines. Although the promoter was repressed in HeLa and PC12 cells, basal levels were 10-30 fold higher in BHK cells. This high basal activity limits the use of the tetracycline-repressed promoter. If a construct could be engineered that was also regulated at the post-transcriptional level, full repression of such a promoter could be achieved in a wider variety of cells. There are several inducible promoters available, however, no one has developed a system to regulate gene expression at the post-transcriptional level. The posttranscriptional regulation of mRNA destabilization of a gene is another possibility for designing regulatable gene expression constructs. If the cDNA can be destabilized by a non-toxic nutrient or analogue, then expression can be stabilized when the analogue is withdrawn.

2B: The invention:

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The cis-acting elements destabilizing mRNA in response to high extracellular glucose were identified by several criteria in the exon encoding the C-terminal 52 amino acids for PKCBII and are shown below. The PKCBII-specific exon is inserted into mature mRNA via alternative splicing of pre-mRNA. The elements inserted may form stem-loop structures providing the secondary structure recognized by destabilizing endonuclease-protein complexes that break A-T bonds (1).

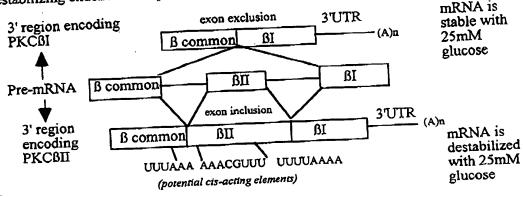


FIGURE 1: Diagram of 3' exons encoding PKCBI and PKCBII via exon inclusion/exclusion and regions of PKCAII exon involved in destabilization of mRNA.

The PKCß gene encodes two isoforms of the serine/threonine kinase that differ by their C-terminal fifty to fifty-two amino acids. If the exon for PKCßII is included in the mature mRNA transcript, PKCßII protein is encoded, the PKCßI exon is not translated due to a stop codon within the exon. If the PKCßII exon is excluded, PKCßI protein is encoded. The 3' untranslated region (UTR) of PKCßI is identical for both PKCßI and PKCßII mRNA. The mRNA for PKCßII is rapidly destabilized in the presence of high extracellular glucose (25 mM) in a number of cells. Like transcription, RNA processing, and translation, mRNA decay is a precise process dependent on specific cis-acting sequences and transacting factors. There are specific pathways triggering mRNA decay including poly (A) shortening, arrest of translation, and endonucleolytic cleavage. We have found that high extracellular glucose concentrations destabilize PKCßII mRNA in a rapid manner, and a nuclease activity in the cytoplasm is involved.

Analysis of the PKCBII exon sequence reveals multiple potential cis-acting elements that may be involved in the destabilization of the PKCBII sequence. These elements may form stem-loop structures that are recognized by putative carbohydrate response-acting factors to target the sequence for decay by cytosolic endonucleases (1).

The sequence could be placed down-stream of the cDNA insert to be expressed in cells, and high extracellular glucose or an analog could be maintained in the media to keep the mRNA of interest down-regulated. When expression of the gene is desired, lowering cell glucose levels to normal (5.5 mM) levels would allow for expression of the target mRNA. Conversely, a gene could be rapidly down-regulated by elevating high extracellular glucose. If the destabilization signal sequence were cloned into a plasmid also carrying the tetracycline repressor, it could fine-tune the system to keep basal levels of the gene low. Glucose or a non-metabolizable analogue provides an inexpensive, non-toxic alternative for regulating gene expression as opposed to expensive and potentially toxic antibiotics.

2C. Detailed description of the invention: The BII-exon should destabilize a cDNA insert in the presence of high extracellular glucose. For our initial sutdies, the pßglobin (pßG) vector (obtained from Norman P. Curthoys, Colorado State University) was used (2). The chimeric phG-BII plasmid was constructed by inserting the PKCBII exon and flanking regions as shown below into the vector at a multicloning site. The vector was created by inserting the PvuII-Bgl II fragment of pSVB10, the portions of the first three exons and the two introns of the rabbit \(\beta\)-globin gene, into the HindIII site of the pRc/RSV vector. The B-globin genomic sequence extends from 9 bp upstream of the transcription initiation site to the translation stop codon. pBG contains a strong viral promoter derived from the long terminal repeat of the Rouse sarcoma virus followed by genomic DNA containing the transcriptional start site, the 3' nontranslated region, the full coding sequence and two introns of the rabbit ß-globin gene, a MCS with four restriction sites, and the 3'nontranslated region and polyadenylation site of the bovine growth hormone gene. Other chimeric constructs will also be tested such as CAT and luciferase to validate the effect of the BII exon (3). The minimal region of the sequence will be used in tetracycline repressor construct systems (4). The chimeric reporter constructs are tested in a number of cell types. We have found that glucose destabilizes PKCBII mRNA in L6 rat skeletal muscle cells, rat aortic vascular smooth muscle cells, and MCF-7 breast cancer cells. Cells such as NIH-3T3 fibroblasts, HeLa cells, PC12 cells, and other tumorigenic and normal cell lines will also be tested for the ability of glucose to destabilize the chimeric construct.

To further define the minimal boundaries of the cis-elements, a portion of the 3' region of the the PKCBII mRNA as diagrammed above was obtained as a 404 bp insert containing the entire 216 bp BII exon and 3' and 5' sequences from the flanking common and BI exon (4). This insert will be restricted by deletion mutagenesis to further limit the amount of mRNA necessary for destabilizing genes of interest.

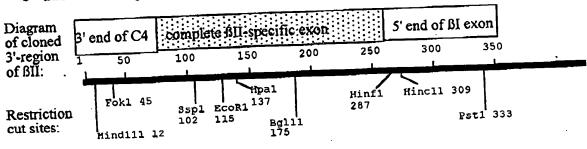
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The destabilization of the PKCBII mRNA by high extracellular glucose is specific for PKCBII mRNA, PKCBI mRNA is not destabilized (4). To determine whether glucose or its metabolites destabilized the mRNA, two glucose analogues were tested. 3-O-methylglucose is taken up by cells but not phosphorylated by hexokinase. 2-Deoxyglucose is taken up by cells and phosphorylated but not further metabolized. Both analogues destabilized PKCBII mRNA. Thus, the further metabolism of glucose to metabolized. Both analogues destabilized PKCBII mRNA. This offers another means of regulating mRNA glucose-6-phosphate is not required for this effect. This offers another means of regulating mRNA stability of the chimeric construct using glucose analogues.

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2D. Drawing of the insert.

Figure 2: Restriction map of 3' region of PKCBII exon with portions of 3'- and 5'-flanking exons to be cloned downstream of the cDNA of interest.



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- 1. Patel, N. A., Chalfant, C.E., Yamamoto, M., Watson, J.E., Eichler, D.C., Cooper, D.R. Acute hyperglycemia regulates transcription and post-transcriptional stability of PKCBII mRNA in vascular smooth muscle cells. FASEB 13, No. 1: 103-113.
- 2. Hansen, W.R., Barsic-Tress, N., Taylor, L., Curthoys, N.P. The 3'-nontranslated region of rat renal glutaminase mRNA contains a pH-responsive stability element. Am. J. Physiol. 271 (Renal Fluid Electrolyte Physiol. 40): F126-F131, 1996.
 - 3. Amara, F.M., Sun, J., Wright J.A. Defining a novel cis-element in the 3'untranslated region of mammalian ribonucleotide reductase component R2 mRNA. J. Biol. Chem. 271: 20126-20131, 1996.
 - 4. Ackland-Berglund, C.E., Leib, D.A., Efficacy of tetracycline-controlled gene expression is influenced by cell type. Biotechniques 18: 196-200, 1995.

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(1) the name of the scientific or professional organization:

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(3) the date of the gathering:

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Tech ID (next no. in sequence)	99A010			
Marketing Status (chosen from drop down list				
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Date on Disclosure	3/11/99			
Date Disclosure received by DPL				
Publication/Presentation Date(s)				
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